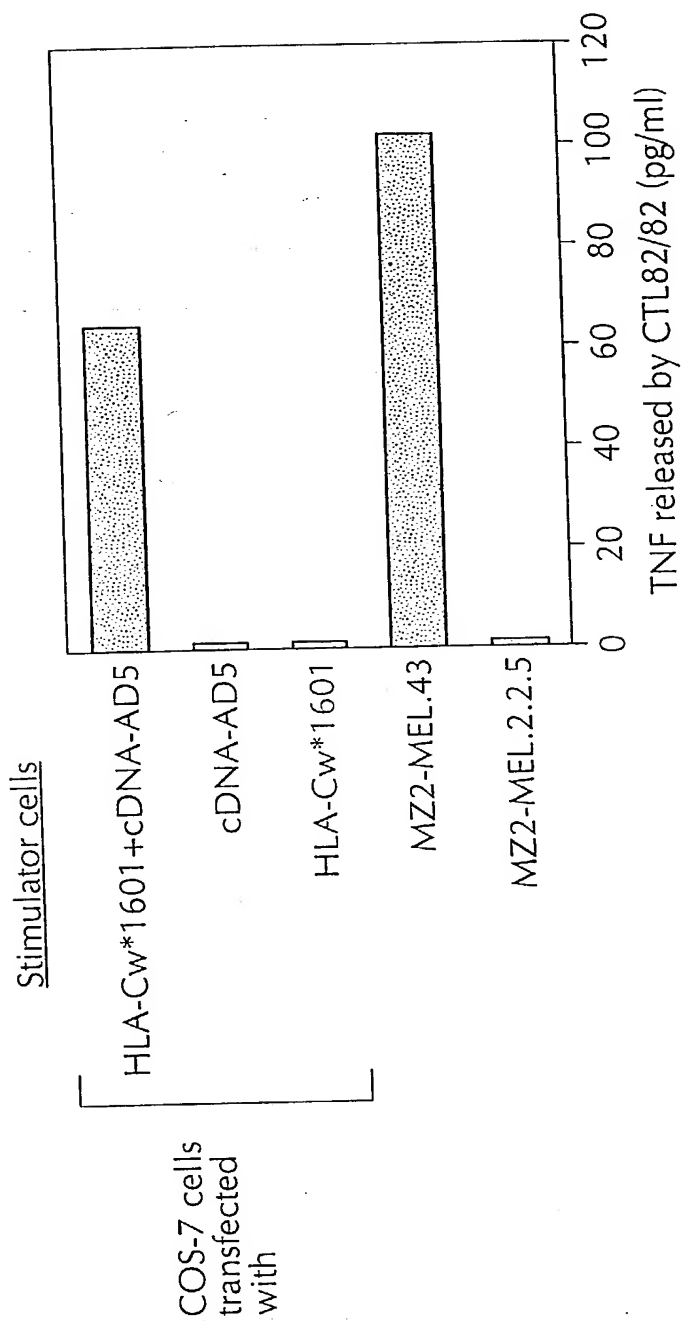


FIG. 2



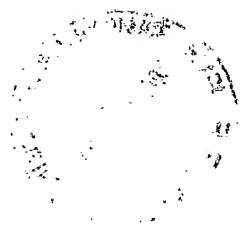


FIG. 3A

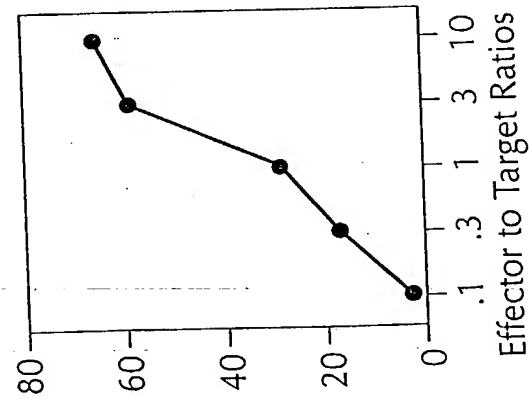


FIG. 3B

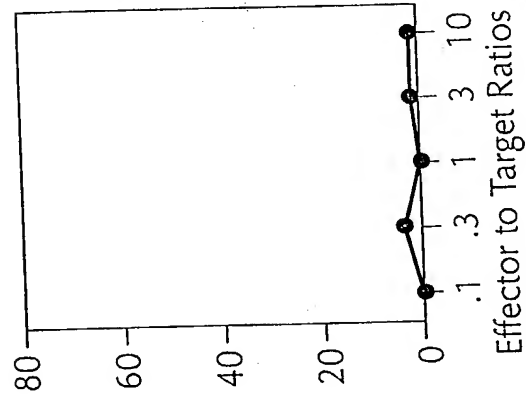


FIG. 3C

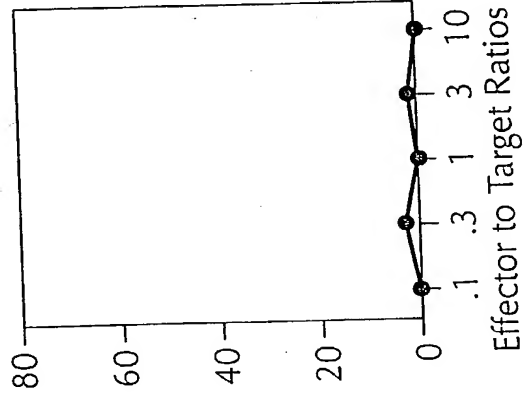


FIG. 3D

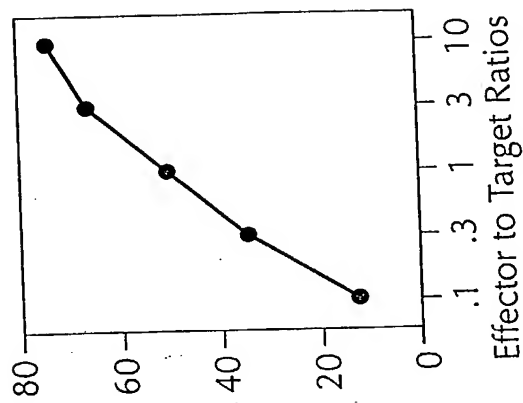


FIG. 4

CGCCAATTTA GGGTCTCCGG TATCTCCCGC TGAGCTGCTC TGTTCCCGGC TTAGAGGACC 60  
 AGGAGAAAGG GGAGCTGGAG GCTGGAGCCT GTAAACACCGT GGCTCGTCTC ACTCTGGATG 120  
 GTGGTGGCAA CAGAGATGGC AGCGCAGCTG GAGTGTTAGG AGGCGGCGCT GAGCGGTAGG 180  
 AGTGGGGCTG GAGCAGTAAG ATGGCGGCCA GAGCGGTTTT TCTGGCATTG S A Q L 240  
 L L Q A R L M K E E S P V V S W R L E P 33  
 TGCTCCAAGC CAGGCTGATG AAGGAGGAGT CCCCTGTGGT GAGCTGGAGG TTGGAGCCTG 300  
 E D G T A L C F I F 43  
 AAGACGGCAC AGCTCTGTGC TTCATCTTCT GAGGTTGTGG CAGCCACGGT GATGGAGACG 360  
 GCAGCTCAAC AGGAGCAATA GGAGGAGATG GAGTTTCACT GTGTCAGCCA GGATGGTCTC 420  
 GATCTCCTGA CCTCGTGATC CGCCCGCCTT GGCCTTCCA AGTGCCGAGA TTACAGCGAT 480  
 GTGCATTTTG TAAGCACTTT GGAGCCACTA TCAAAATGCTG TGAAGAGAAA TGTACCCAGA 540  
 TGTATCATTA TCCTTGTGCT GCAGGAGCCG GCTCCTTTCA GGATTTCACT CACATCTTCC 600  
 TGCTTTGTCC AGAACACATT GACCAAGCTC CTGAAAGATG TAAAGTTTACT ACGCATAGAC 660  
 TTTTAAACTT CAACCAATGT ATTTACTGAA AATAACAAAT GTTGTAATTT CCTGTAGTGT 730  
 TATTCTACTT GTATTAAAAG GTAATAATAC ATAATCATTA AAATCTGAGG GATCATTGCC 780  
 AGAGATTGTT GGGAGGGGAA ATGTTATCAA CGGTTTCACT GAAATTAAT CCAGAAAAGTT 840  
 ATTTCCCTCAG AAAAATCAAA TAAAGTTTGC ATGTTTTTTA TTCTTAAAC ATTTTAAAAA 900  
 CCACTGTAGA ATGATGTAAA TAGGGACTGT GCAGTATTTT TGACATATAC TATAAAATTA 960  
 TTAAAAAGTC AATCAGTATT CAACATCTTT TACACTAAAA AGCC 1004

1000011000.5454000

FIG. 5

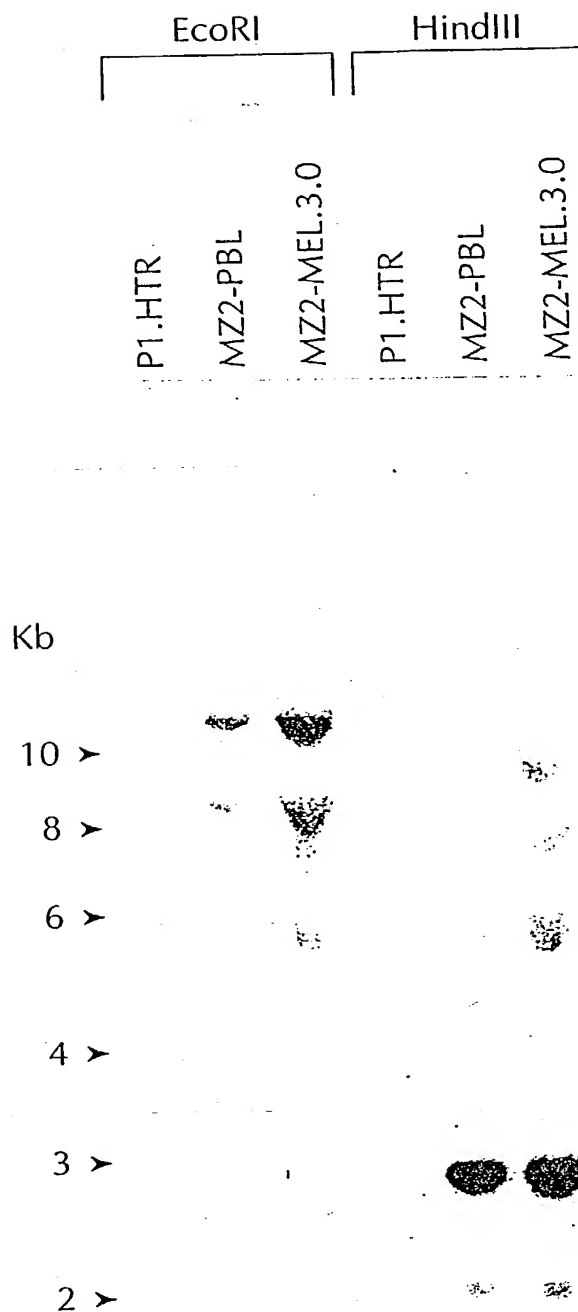


FIG. 6

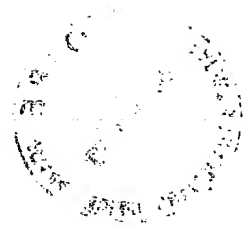
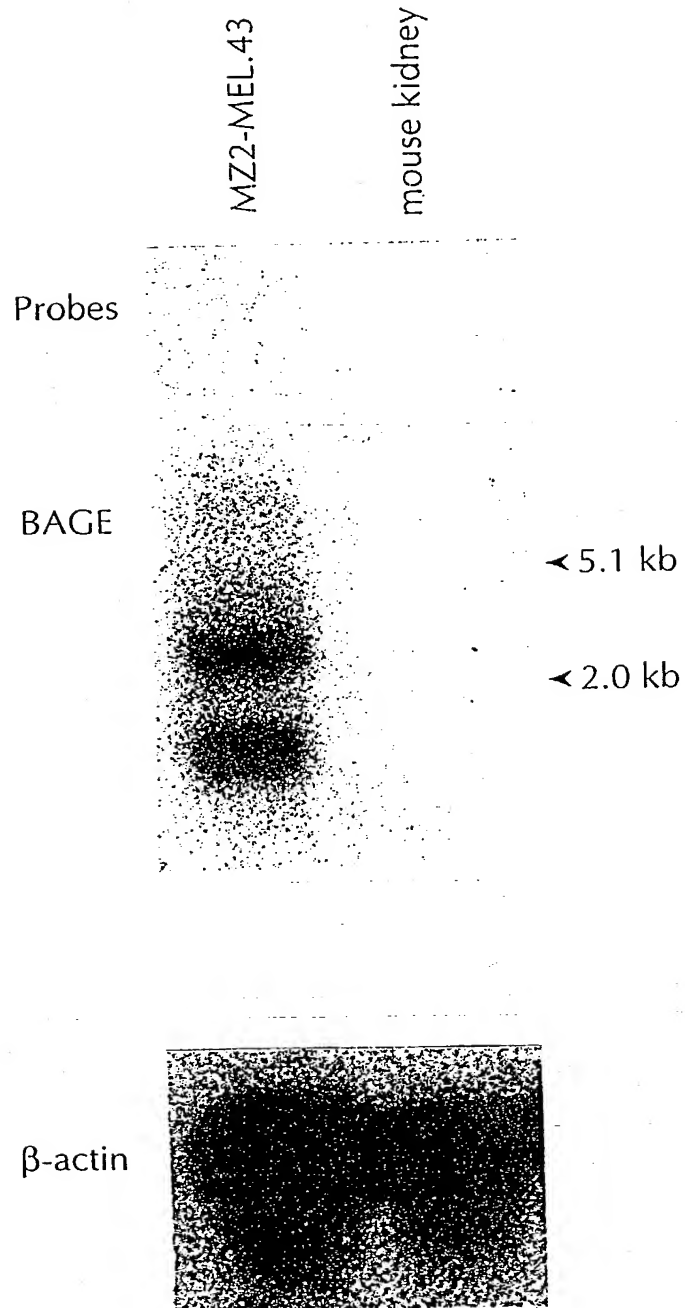


FIG. 7

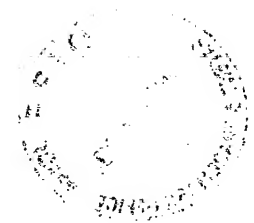
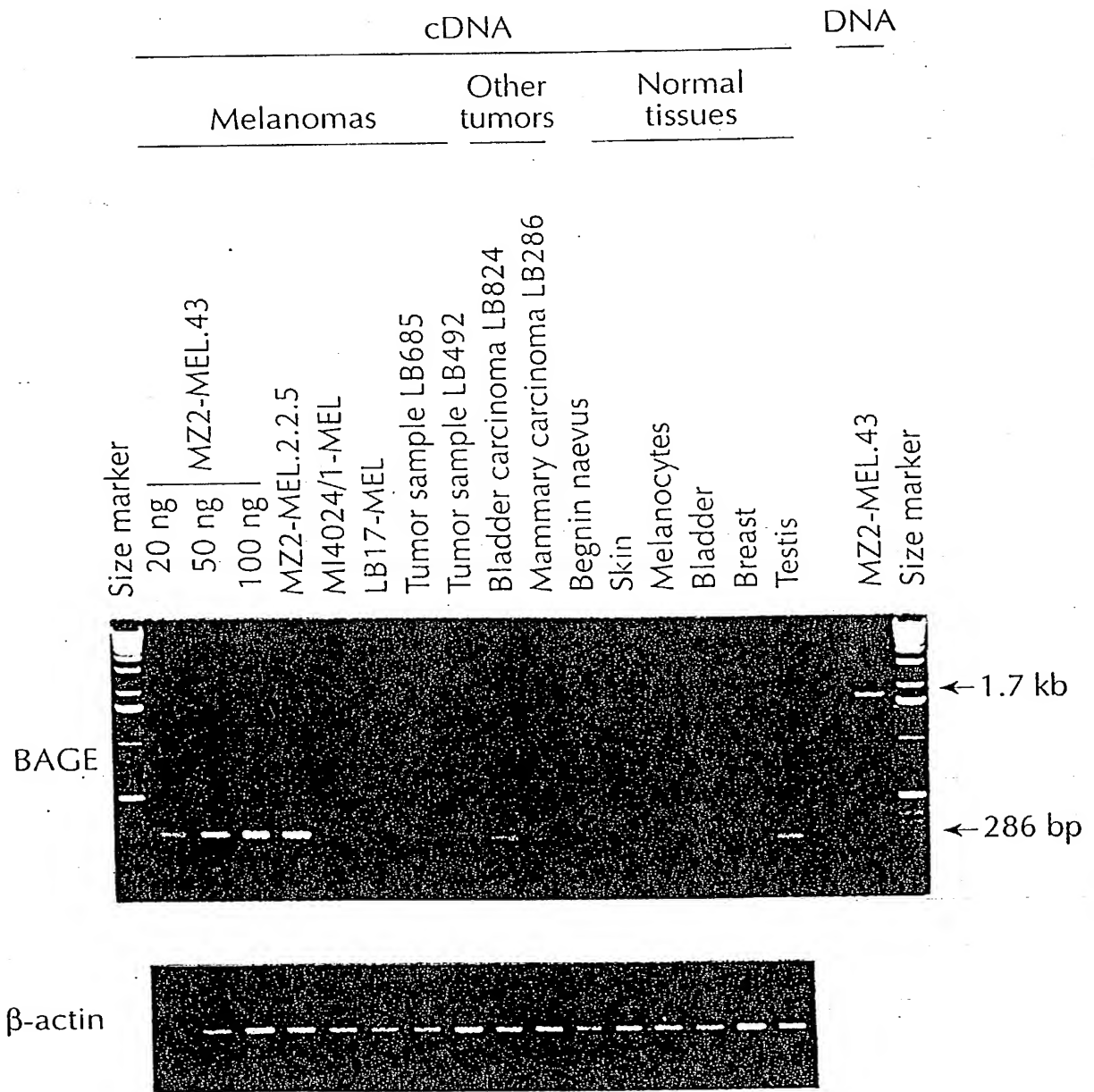


FIG. 8

